==========

Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: Mon Jun 04 19:15:40 EDT 2007

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****

Reviewer Comments:

<110> ARES TRADING S.A.
<110> FAGAN, Richard Joseph
<110> DAVIDS, Andrew Robert
<110> PHELPS, Christopher Benjamin
<110> POWER, Christine
<110> BOSCHERT, Ursula
<110> CHVATCHKO, Yolande

Per 1.823 of the Sequence Rules, the <110> numeric identifier is only shown on the first applicant's line; please delete the additional <110>'s.

<140> PCT/GB2004/004772
<141> 2004-11-12

Please change the <140> to <150> and the <141> to <151>, since these are prior application data. They are not the current application number and current filing date.

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Validated By CRFValidator v 1.0.2

Application No: 10579113 Version No: 1.0

Input Set:

Output Set:

Started: 2007-05-22 13:27:47.914
Finished: 2007-05-22 13:27:48.874
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 960 ms
Total Warnings: 0
Total Errors: 7
No. of SeqIDs Defined: 31
Actual SeqID Count: 31

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E 249	Order Sequence Error <110> -> <110>; Expected Mandatory Tag: <120> in Header
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E 250	Structural Validation Error; Sequence listing may not be indexable

SEQUENCE LISTING

<110> ARES TRADING S.A.
<110> FAGAN, Richard Joseph
<110> DAVIDS, Andrew Robert
<110> PHELPS, Christopher Benjamin
<110> POWER, Christine
<110> BOSCHERT, Ursula
<110> CHVATCHKO, Yolande

<120> CYTOKINE AGONIST MOLECULES

<130> P035815WO

<140> PCT/GB2004/004772
<141> 2004-11-12

<150> GB0326393.6
<151> 2003-11-12

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<211> 29
<212> PRT
<213> Homo sapiens

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Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp
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<212> DNA
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gcaccctgcg gcctgactat cgagaccgta tccgactctt tgaaaatggc tccctgcttc 240
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<210> 4
<211> 114
<212> PRT
<213> Homo sapiens

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Thr Val Gly Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser
20 25 30

Ser Asp Arg Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val
35 40 45

Thr Val Val Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro
50 55 60

Asp Tyr Arg Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu
65 70 75 80

Ser Asp Leu Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser
85 90 95

Ile Thr Asp Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val
100 105 110

Asp Val

<210> 5
<211> 282
<212> DNA
<213> Homo sapiens

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<210> 6
<211> 94
<212> PRT
<213> Homo sapiens

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20 25 30

Pro Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser
35 40 45

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Arg Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val
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<213> Homo sapiens

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<212> PRT
<213> Homo sapiens

<400> 8

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<211> 74
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<213> Homo sapiens

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<210> 10
<211> 25
<212> PRT
<213> Homo sapiens

<400> 10

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Asn Asp Asp Arg Leu Lys Pro Glu Ala
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<211> 71
<212> DNA
<213> Homo sapiens

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<210> 12
<211> 23
<212> PRT
<213> Homo sapiens

<400> 12
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Leu Tyr Ile Leu Lys Asp Lys
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<210> 13
<211> 303
<212> DNA
<213> Homo sapiens

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<210> 14
<211> 100
<212> PRT
<213> Homo sapiens

<400> 14
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Arg Ser Pro Gly Leu Pro Ile Arg Ser Ala Arg Arg Tyr Pro Arg Ser
35 40 45

Pro Ala Arg Ser Pro Ala Thr Gly Arg Thr His Ser Ser Pro Pro Arg
50 55 60

Ala Pro Ser Ser Pro Gly Arg Ser Arg Ser Ala Ser Arg Thr Leu Arg
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Thr Ala Gly Val His Ile Ile Arg Glu Gln Asp Glu Ala Gly Pro Val
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Glu Ile Ser Ala
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<210> 15

<211> 1251
<212> DNA
<213> Homo sapiens

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<210> 16
<211> 416
<212> PRT
<213> Homo sapiens

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20 25 30

Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly
35 40 45

Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg
50 55 60

Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val
65 70 75 80

Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg
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Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu
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Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp
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Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro
                165                 170                 175

Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg
            180                 185                 190

Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu
        195                 200                 205

Met Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser
    210                 215                 220

Gln Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser
225                 230                 235                 240

Leu Tyr Ile Ile Leu Ser Thr Gly Gly Ile Phe Leu Leu Val Thr Leu
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Val Thr Val Cys Ala Cys Trp Lys Pro Ser Lys Arg Lys Gln Lys Lys
            260                 265                 270

Leu Glu Lys Gln Asn Ser Leu Glu Tyr Met Asp Gln Asn Asp Asp Arg
        275                 280                 285

Leu Lys Pro Glu Ala Asp Thr Leu Pro Arg Ser Gly Glu Gln Glu Arg
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Lys Asn Pro Met Ala Leu Tyr Ile Leu Lys Asp Lys Asp Ser Pro Glu
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Thr Glu Glu Asn Pro Ala Pro Glu Pro Arg Ser Ala Thr Glu Pro Gly
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Pro Pro Gly Tyr Ser Val Ser Pro Ala Val Pro Gly Arg Ser Pro Gly
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Leu Pro Ile Arg Ser Ala Arg Arg Tyr Pro Arg Ser Pro Ala Arg Ser
        355                 360                 365

Pro Ala Thr Gly Arg Thr His Ser Ser Pro Pro Arg Ala Pro Ser Ser
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Pro Gly Arg Ser Arg Ser Ala Ser Arg Thr Leu Arg Thr Ala Gly Val
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<211> 1257

<212> DNA
<213> Mus musculus

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<211> 418
<212> PRT
<213> Mus musculus

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Leu Ser Pro Phe Val Tyr Leu Leu Leu Ile Gln Pro Val Pro Leu Glu
20 25 30

Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly
35 40 45

Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Lys
50 55 60

Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val
65 70 75 80

Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg
85 90 95

Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu
100 105 110

Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp
115 120 125

Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro

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Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu
145                 150                 155                 160

Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro
                165                 170                 175

Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg
            180                 185                 190

Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu
        195                 200                 205

Met Glu Asp Asp Asp Leu Tyr Ser Cys Val Val Glu Asn Pro Ile Ser
    210                 215                 220

Gln Val Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser
225                 230                 235                 240

Leu Tyr Ile Ile Leu Ser Thr Gly Gly Ile Phe Leu Leu Val Thr Leu
                245                 250                 255

Val Thr Val Cys Ala Cys Trp Lys Pro Ser Lys Lys Ser Arg Lys Lys
            260                 265                 270

Arg Lys Leu Glu Lys Gln Asn Ser Leu Glu Tyr Met Asp Gln Asn Asp
        275                 280                 285

Asp Arg Leu Lys Ser Glu Ala Asp Thr Leu Pro Arg Ser Gly Glu Gln
    290                 295                 300

Glu Arg Lys Asn Pro Met Ala Leu Tyr Ile Leu Lys Asp Lys Asp Ser
305                 310                 315                 320

Ser Glu Pro Asp Glu Asn Pro Ala Thr Glu Pro Arg Ser Thr Thr Glu
                325                 330                 335

Pro Gly Pro Pro Gly Tyr Ser Val Ser Pro Pro Val Pro Gly Arg Ser
            340                 345                 350

Pro Gly Leu Pro Ile Arg Ser Ala Arg Arg Tyr Pro Arg Ser Pro Ala
        355                 360                 365

Arg Ser Pro Ala Thr Gly Arg Thr His Thr Ser Pro Pro Arg Ala Pro
    370                 375                 380

Ser Ser Pro Gly Arg Ser Arg Ser Ser Ser Arg Ser Leu Arg Thr Ala
385                 390                 395                 400

Gly Val Gln Arg Ile Arg Glu Gln Asp Glu Ser Gly Gln Val Glu Ile
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Ser Ala
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<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu
            20   25   30